

FIG. 1

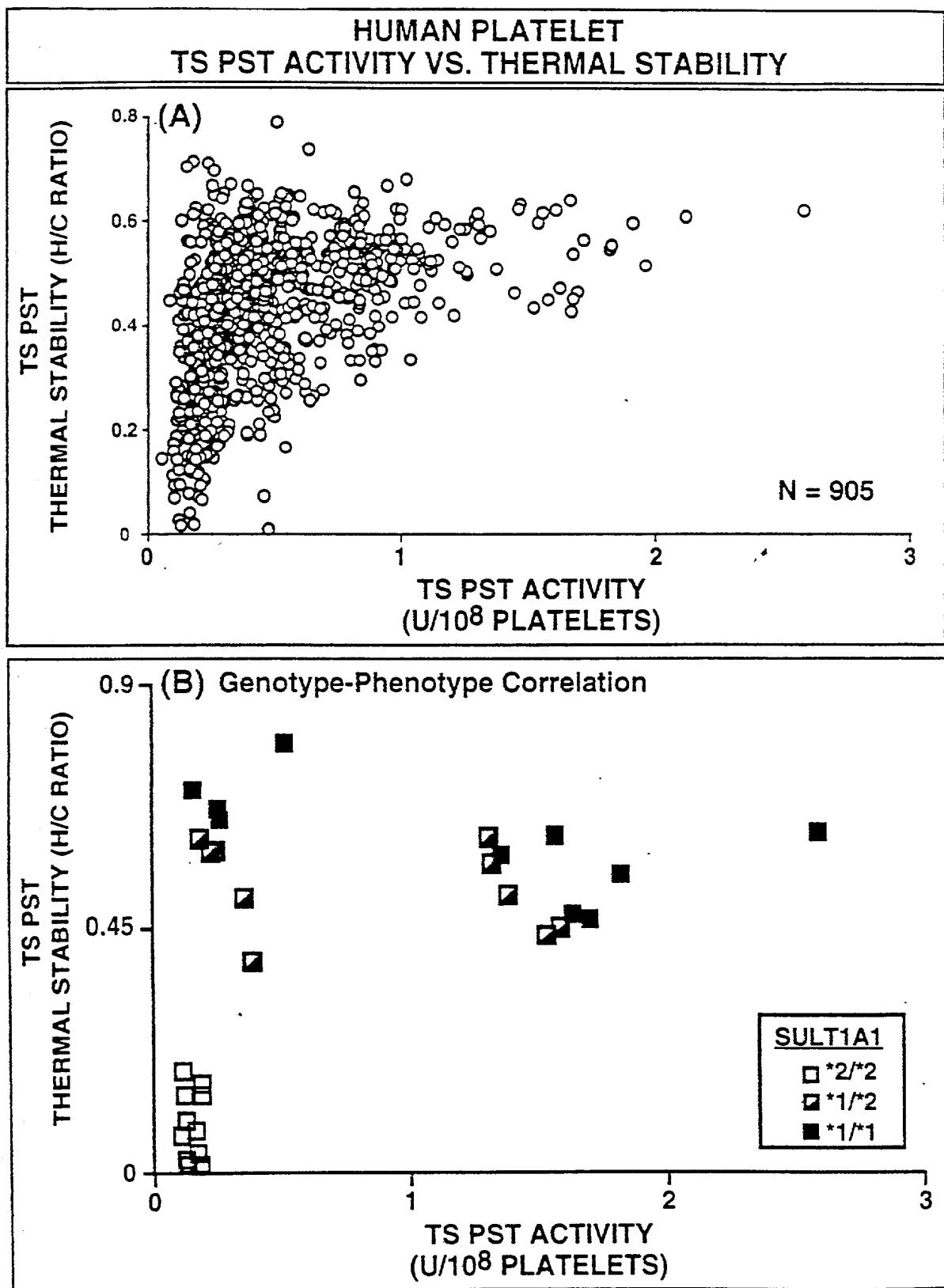


FIG. 2

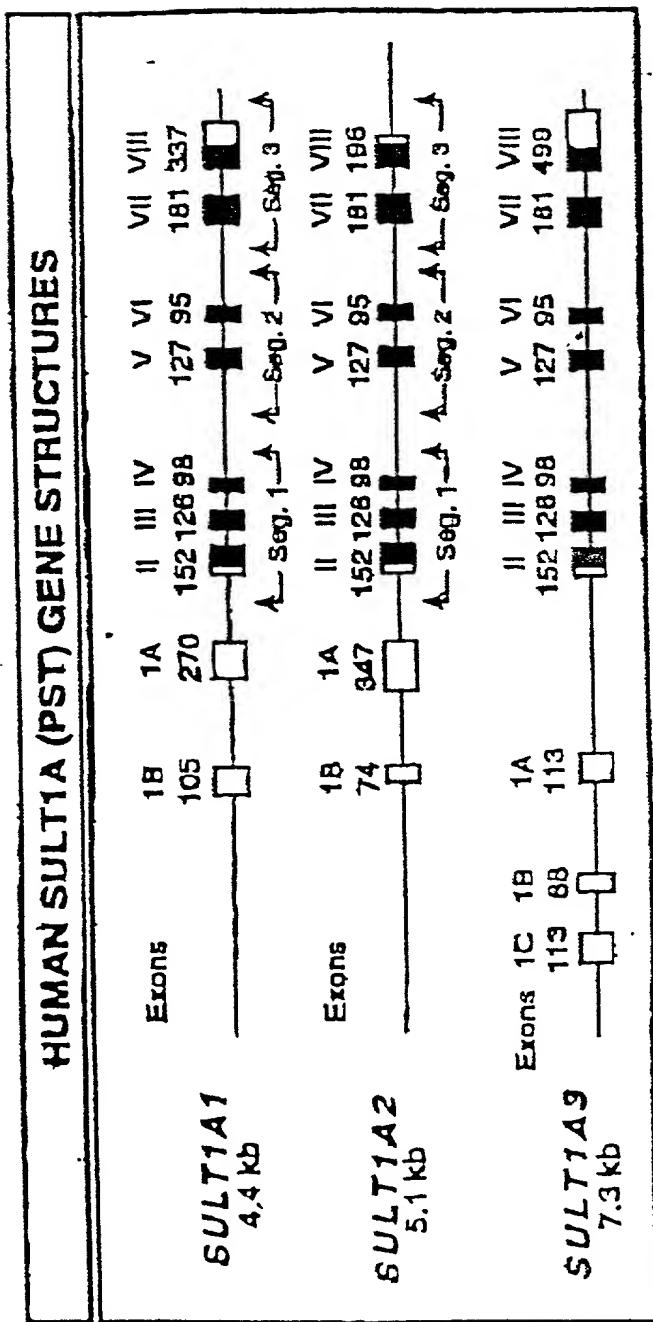


FIG. 3

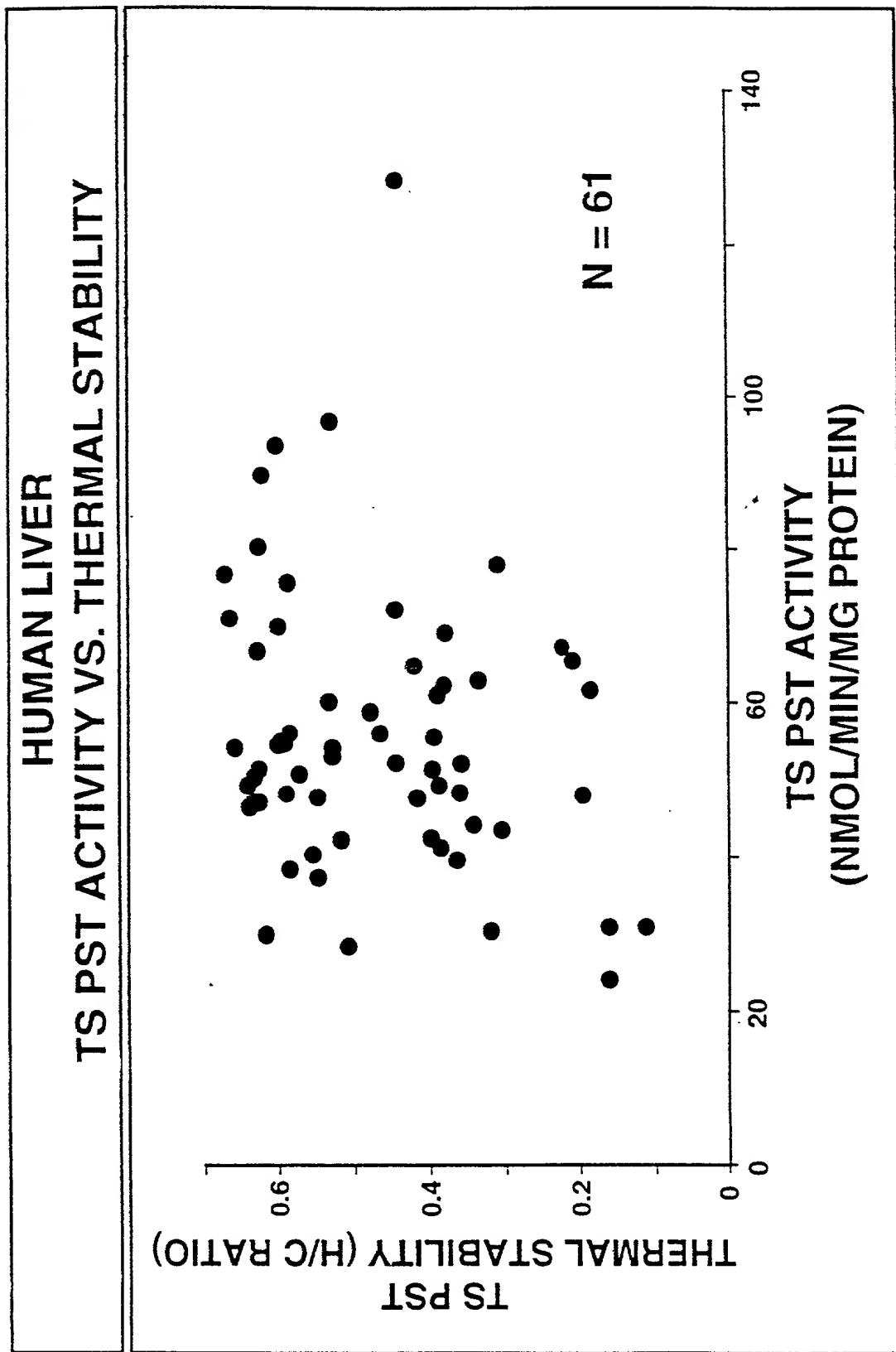


FIG. 4

HUMAN LIVER
TS PST GENOTYPE-PHENOTYPE CORRELATION

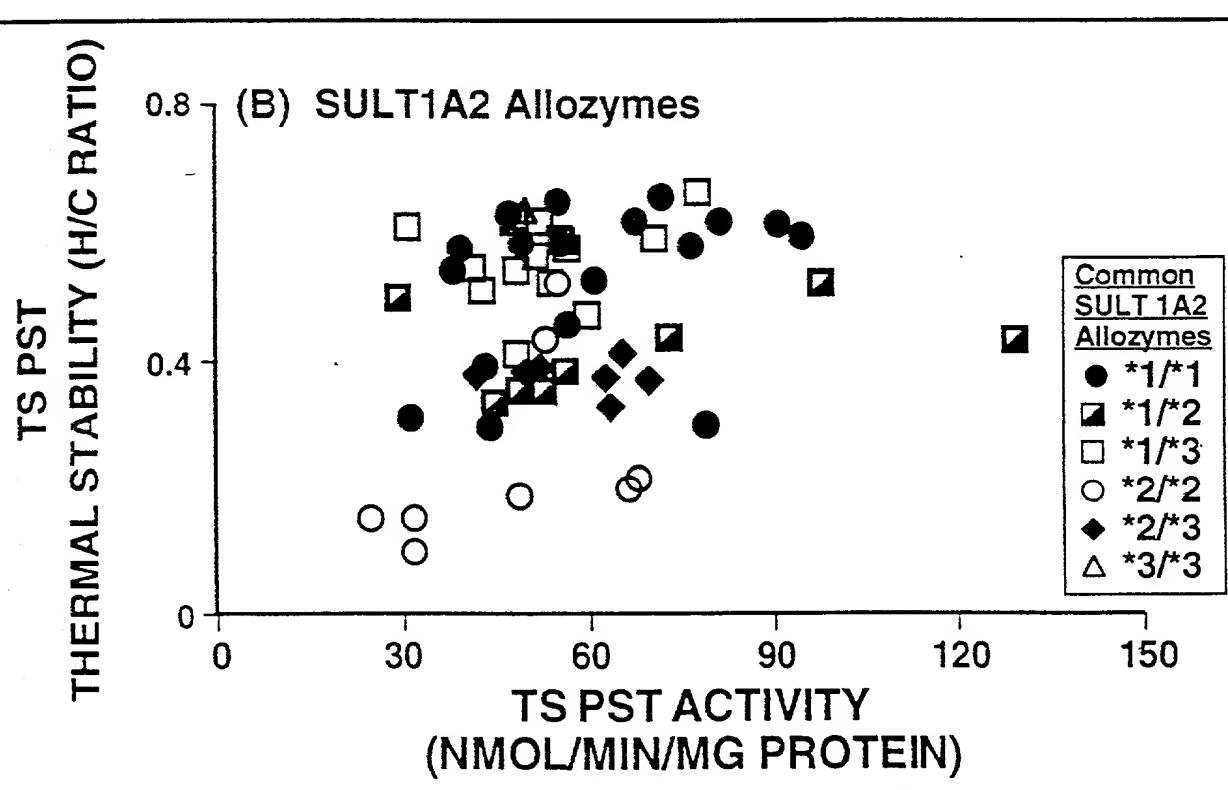
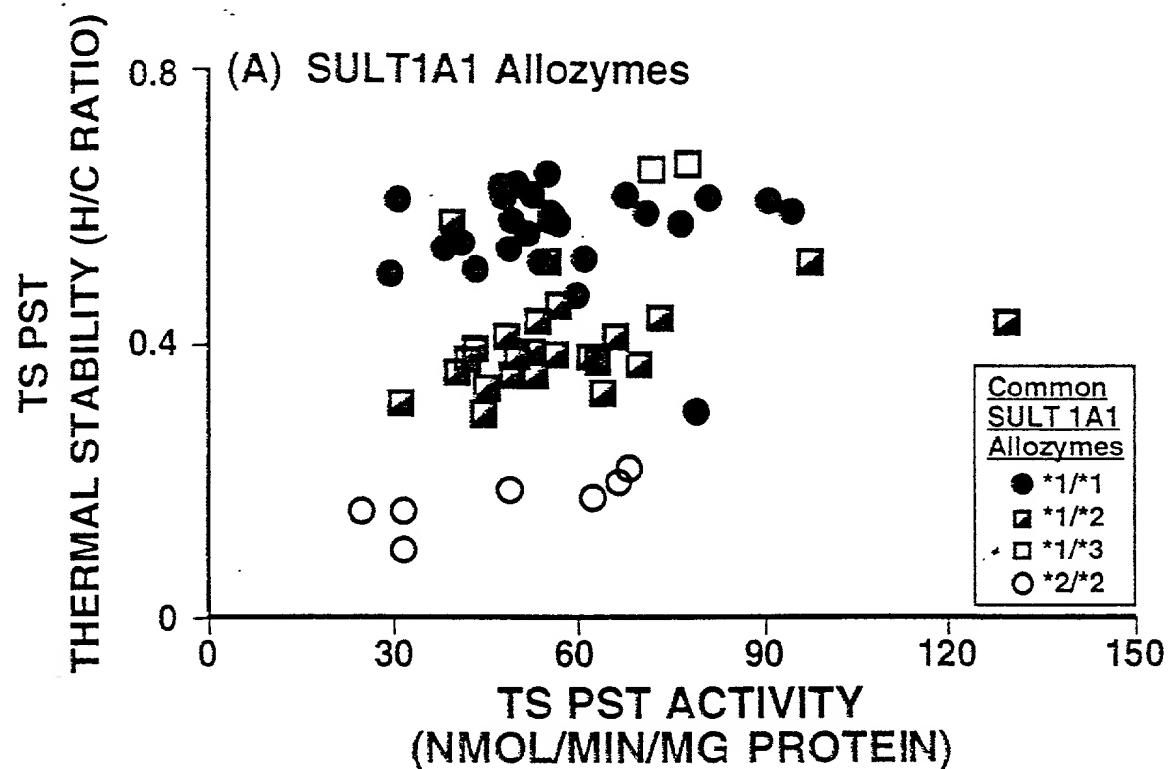


FIG. 5

HUMAN SULT1A1 GENE SEQUENCE

Figure 1. *SOX11A1* gene sequence. Exon sequence is shown in uppercase, while intron, 5'-flanking region and 3'-flanking region sequence is in lowercase. The 'ATG' translation initiation codon and the 'TGA' translation termination codon are underlined. Amino acid sequence encoded by the gene is listed in single-letter code beneath the nucleotide sequence.

FIG. 6

HUMAN SULT1A2 GENE SEQUENCE

Figure 2 *Sty222* gene sequence. Exon sequence is shown in uppercase, while intron, 5'-flanking region and 3'-flanking region sequence is in lowercase. Amino acid sequence encoded by the gene is listed in single-letter code beneath the nucleotide sequence.

FIG. 7

FIG. 7 CONT.

3241 ctgagtagct ggggttacaa gagcaagcca ccacacccag ctaattttga aaaatatttt
 3301 tttagagga gaggtcttgc ttgttgtcc aggttggct caaactccag ggctcaaggg
 3361 atcccttccc gttggctcc caaggctctg ggattacagg cgggagtcac cctgcctggg
 3421 ccccttctt ttagttagtca tcagtttca ttcccgacag aggctctagc ccctggtacc
 3481 agcttagttg ctcaatggc tttgtttgtt ctggagccca gatggactgt ggccaggcaa
 3541 gtggatcaca gacctggccg gcctgggagg ttccacatg tgagggggcat gaggggggct
 3601 caaggagggg agcatcgggg agaggagcgc actgggtgga ggctgggggt cccagcagga
 3661 aatggtgaga caaaggcgc tggctggcag ggagacagca caggcaggcc ctagagcttc
 3721 ctcagcacag ctggacttc ctggagaccc tcacacaccc tgatatctgg gccccgcgt
 3781 acgaggggtgc tttcaactgt ctgcactatg cccaggccc tgggattttg aacagctctg
 3841 caggtgactg aaaggtgcgg ccaggctgg gaacgacctg gttcagccc cagccccgccc
 3901 actgactgac tttgtgatgt cgggcaagtc actcagccctc ctagggcttc agtacttcc
 3961 ctgaaaagcaa aaactctgca aaggggcagc tgggtgctgg ctcacaccc tgatccac
 4021 actttggag gctgaggtag acaaatact tgaggccagg agttctagac cagcctggcc
 4081 aacatggtga aacccatct ctactaaaga aaaaaaaaaa ttagtgcagc atgggtgtac
 4141 atgcttgcata tccagctac tgggatgcc gaggcgggag gattgcttga acccaagagg
 4201 tggagttgc agtgcactgatgt gattgtgcca cactgcactc cagctgggt gagagtgaga
 4261 ctccatctca aaaaaaaaaa aaaaaagaga gaatcccact ttcttgctgt tggatggtg
 4321 gtaaggaaac gggctggct ctggccctg atgcacggaa atggagctga tccaggacac
 4381 ctcccccccg ccactggagt acgtgaaggg ggtccgcgtc atcaagtact ttgcagaggc
 4441 actggggccc ctgcacagct tccaaaggccg acctgtatgc ctgctcatca acacctaccc
 4501 caagtctgt aagtgaggag gcccacccac cctctccag gccgcgttcc ccacccctgg
 4561 cagcaaggcgtc gtgcctcag cctgctcacc tcctatctcc ctcctctcc aggccaccacc
 4621 tgggtgagcc agataactgga catgtatctac cagggcggcg acctagaagaa gtgtaaacccgg
 4681 gctcccatct acgtacgggt gcccctcctt gaggtcaatg atccagggga accctcaggt
 4741 gcatggctgg gtcctggggg taagggaaatgggaaagac agggctgggg cttcagctca
 4801 ccagacccctc cctgacccac tactcaggc tggagactct gaaagacaca ccccccac
 4861 ggctcatcaa gtcacacccctg cccctggctc tgctccctca gactcttttgc gatcagaagg
 4921 tcaaggtag gcccggctca atgggtcaca cctgtcatcc cagtttgaga ctgaggaggg
 4981 aggtccctt gaaggcgaga gatggagacc agcctggca acattgtgt agagatgaca
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 5161 tgatcctgt gtcactcc acgctggca acacagcaaa accatctta cggaaaaaaa
 5221 agttccact gactggcaag gaaagccagg aaggggggct caggtgcct ctcagccatg
 5281 tacctgttct tctggaaaggg ctcctcgct tctgccagc tcataccatc tttttttt
 5341 ttgagacaga gtcttgctt gtcacccctgg ctggagtgca gtggcatgtat ctcagctcac
 5401 tgcaacccctc gcccctccag ttcaagtgtat tctcctgcct cagcctcctg agtagctgg
 5461 attacaggcc tggatccatca caccggcata atttttgtat tcttttttagt agagacgggg
 5521 tttcaccatg tggtaagt ggtatctcaaa ctcttgaccc tggatctc tggctcgac
 5581 ctcacaaaatg gtcggaaatc caggcgtgag ccaccgcgccc tggcccttt ttttttttag
 5641 acagtttccac ttttggcc gaggctagag cgcaatcgtg tgatctcggt tcactgcaac
 5701 caccggcctcc tgggttcaag caattctcc tggatctc cccaaaggagc tggattaca
 5761 ggtacctgccc accacggcccg gctaattttg tatttttagt agagatgggg tttcaccatg
 5821 ttgggtcaggc tgggttgcata ctcctgcaccc caggtgatct ggcacccctgg ccccccgg
 5881 tgccgggatt agaggcatga gcccacccgc ccagcctca tcacatctt agagaggaca
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 6181 gtttggccctt acctgtgcacg tgagatctca agcaagtcac cctctaaagcc tcagggttcc
 6241 cttttggccat ttcaacagat gagctggctt ggggtgggtt gtgtgggtat ggtgtgggg
 6301 ctgggtccctc tgccctgcata ggtggatctat gttcccgaa acccaaaaggaa cgtggcggtc
 6361 tcctactacc atttccaccg tatggaaaag ggcacccctg agcctgggac ctggggacagc
 6421 ttccctggaaa agttcatggc tggagaaatgg gggcttgcact ggagggaaaggaa ggggtgtgaag
 6481 ccgagggggtg tgggttataa cgtacagccaa ccctgtgtcg gtggccctg cccgcttctc
 6541 tagtgcctca cgggtccctgg taccagcagc tgcaaggatgt gtgggagctg agccgcaccc
 6601 accctgttct ctcaccccttc tatgaagata tgaaggaggt gagaccgact gtgtatgcctc

FIG. 7 CONT.

6661 cccccatgtg acacctgggg gcaggcacct cacagggacc caccaaggcc acccagccc
6721 gtcctgggc ggctcccaca gcaagccgg attcccccac tcacccctt ggcggcaggcc
6781 cccccactgc agccccacct ggcagcaggc tcggcacagc tttcatctc tgcacctgag
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7021 gggccccagg ggtttaggct ttgtgggtt tctagggaa gtggccagat cccctctgag
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7381 gcaacttgggg cctccctgg gatgagactc cagctttgtt ccctgccttc ctccccccagg
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8281 gaggtgtctt caaacaggat agtatagtcc ttccatggggg ttgtgggggtt aggtggccaa
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